STIC Biotechnology Systems Branch

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Application Serial Number: 10/530,/06
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Date Processed by STIC: 4//2/05

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
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 U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street.
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Revised 01/24/05



RAW SEQUENCE LISTING DATE: 04/12/2005 PATENT APPLICATION: US/10/530,106 TIME: 10:20:31

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\04122005\J530106.raw

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     Walchli, Sebastien
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rention and/or
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 14 <130> FILE REFERENCE: SLII-P01-003
> 17 <140> CURRENT APPLICATION NUMBER: US/10/530,106
 17 <141> CURRENT FILING DATE: 2005-04-01
 17 <160> NUMBER OF SEQ ID NOS: 34
 21 <170> SOFTWARE: PatentIn version 3.1
 25 <210> SEQ ID NO: 1
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215

pp 6-7

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/530,106

DATE: 04/12/2005 TIME: 10:20:31

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160	× 7 -	D	~ 1-	~ 1	485	01	03		a	490	~		_		495	
163 A	мта	PIO	GIII	500	PIO	GIY	GIn	ser		Tyr	ser	ıyr	Trp		Ser	Trp
		7 ~~	C1.,		Mot	Th.	7	Daia	505	TD b	a 1		ml	510	~ 1	m1
167 V 168	vaı	Arg	515	GIY	Mec.	1111	Asp		Arg	inr	GIN	ser		ser	GIY	inr
171 2	A cn	Tla		T.011	Lve	Glu	T 011	520	ת דת	C1	C-~	T 011	525	17-1 ~	T	ml
172	rsp	530	1111	neu	ыys	GIU	535	GIU	Ата	GIY	ser		TYL	HIS	Leu	Inr
175: V	(7 - 1		775	Glu	7~~	λαπ		370 J	7~~	~ 1	m	540	C	(T)	T	m\
176 5		пр	AIG	Giu	Arg	550	Giu	val	Arg	GIY		ASI	ser	THE	Leu	
		715	Thr	777	Dro		C1	37-3	The sec	7	555	~1 - -		G3	ml	560
179 <i>1</i> 180	a	nia	TIIT	vra	565	ven	GIU	val	TIIT	570	ьец	GIII	ASI	GIU		
	Th~	Lara	7.00	e o ~		Mot	T 011	TI-v-	m		77-	Desc	a 1	3	575	
183 7 184	- 111	Ly S	von	580	Val	MEC	neu	11D	585	пув	WIG	PLO	GIĀ		PLO	nis
187 8	Ser	Glr	T.e.r		\1=1	Тчг	Trn	7727		Trans	ת דת	C^~	T	590	11i-	Desc
188	-er	3111	595	TAT	val	TYL	тър	600	GIII	тър	ATG	ser		GTĀ	uis	PEO
191 7	مدم	Δrσ		Gl n	Δαρ	Dro	G1 ~		700	Ф~~	17 - T	7 a~	605	The se	C	7
192	9	610	GTĀ	GIII	veħ	FIO	615		ASII	пр	val		·GIN	Inr	ser	Arg
124		910					012	•				620				

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208	207	Cys	Val	Ser	Thr	Ser	Ala	Gly	Tyr	Gly	Val	Asn	Leu	Ile	Trp	Ser	Cys
215 Gly Ser Gln Asp Arg Ser Ser Cys Gly Glu Ala Val Ser Val Leu Gly To To To To To To To T	208			675					680					685			
215 Gly Ser Gln Asp Arg Ser Ser Cys Gly Glu Ala Vai Ser Vai Leu Gly 720 720 720 720 720 735 730 735	211	Pro	Gln	Gly	Gly	Tyr	Glu	Ala	Phe	Glu	Leu	Glu	Val	Gly	Gly	Gln	Arg
216 705 710 715 720 720 720 721 720 720 721 720 722 720 725									•								
Leu Gly Pro Ala Arg Ser Tyr Pro Ala Thr Ile Thr Thr Ile Trp Asp			Ser	Gln	Asp	Arg		Ser	Cys	Gly	Glu		Val	Ser	Val	Leu	_
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232					_			~.		_	_	_	_		_	_	
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236 785 790 795 800 239 Pro Ala Glu Asp Phe Ala Asp Phe Ala Asp Phe 805 815 810 815 815 240 805 810 810 815 815 815 815 243 Asn Cys Gly Phe Ala Asp Glu Tyr Gln Gln Leu Ser Leu Val Gly His 820 825 830 845 830 845 830 845 830 845 830 845 830 845 830 845 830 845 830 845 830 845 830 845 845 830 845 860 845 860 845 855 860 845 860 855 860 855 860 865 860 865 860 865 860 865 860 865 860 865 860 865 860 865 860 865 <td></td> <td>C1 n</td> <td></td> <td>Dwa</td> <td>C1.,</td> <td>T 011</td> <td>7</td> <td></td> <td>T</td> <td>77-7</td> <td>Db -</td> <td>0</td> <td></td> <td>D</td> <td>61</td> <td>3</td> <td>-7 -</td>		C1 n		Dwa	C1.,	T 011	7		T	77-7	Db -	0		D	61	3	- 7 -
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240			Δla	Glu	Acn	Dho		Λcn	uic	17-a T	7.50		N an	C1.,	7. ~~~	7 ~~	
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275 Leu Arg Val Thr Leu Val Gly Glu Glu Val Met Glu Asn Trp Thr Val 276 945 950 955 960 279 Arg Glu Leu Leu Leu Gln Val Glu Glu Glu Lys Thr Leu Ser Val 280 965 970 975 283 Arg Gln Phe His Tyr Gln Ala Trp Pro Asp His Gly Val Pro Ser Ser 284 980 985 990 287 Pro Asp Thr Leu Leu Ala Phe Trp Arg Met Leu Arg Gln Trp Leu Asp 288 995 1000 1005		Cys		птр	IYI	TIP	PIO		Asp	ser	GIII	PIO	_	THE	HIS	GIA	HIS
276 945 950 955 960 279 Arg Glu Leu Leu Leu Leu Gln Val Glu Glu Glu Lys Thr Leu Ser Val 280 965 970 975 283 Arg Gln Phe His Tyr Gln Ala Trp Pro Asp His Gly Val Pro Ser Ser 284 980 985 990 287 Pro Asp Thr Leu Leu Ala Phe Trp Arg Met Leu Arg Gln Trp Leu Asp 1005		T.e.u		Val.	Thr	T.e.11	V=1		Glu	Glu	17a 1	Mot		N en	т~~	Thr	17-1
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280 965 970 975 283 Arg Gln Phe His Tyr Gln Ala Trp Pro Asp His Gly Val Pro Ser Ser 980 985 990 284 Pro Asp Pro Asp Thr Leu Leu Ala Phe Trp Arg Met Leu Arg Gln Trp Leu Asp 1005 1005			Glu	Len	Len	Len		Gln	Val	Glu	Glu		Larg	Thr	T.011	Sar	
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284 980 985 990 287 Pro Asp Thr Leu Leu Ala Phe Trp Arg Met Leu Arg Gln Trp Leu Asp 288 995 1000 1005		Arq	Gln	Phe	His			Ala	Trp	Pro		His	Glv	Val	Pro		Ser
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PATENT APPLICATION: US/10/530,106

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419	<213> ORGANISM: Artificial Sequence		
423	<220> FEATURE:		
425	<223> OTHER INFORMATION: Primer		
427	<400> SEQUENCE: 7	•	
428	catgctgacc aactgcatgg	•	20
	<210> SEQ ID NO: 8		
	<211> LENGTH: 22		
	<212> TYPE: DNA		
437	<213> ORGANISM: Artificial Sequence	•	
	<220> FEATURE:	·	
	<223> OTHER INFORMATION: Primer		
	<400> SEQUENCE: 8		
	gatgggattt ccattgatga ca	•	22
	<210> SEQ ID NO: 9		
	<211> LENGTH: 18		
	<212> TYPE: DNA		
	<213> ORGANISM: Artificial Sequence		
	<220> FEATURE:	,	
	<223> OTHER INFORMATION: Primer		
	<400> SEQUENCE: 9	•	
	ccacccatgg caaattcc	·	18
	<210> SEQ ID NO: 10	•	
	<211> LENGTH: 21		
	<212> TYPE: DNA		
	<213> ORGANISM: Artificial Sequence		
	<220> FEATURE:		
	<223> OTHER INFORMATION: Primer		
	<400> SEQUENCE: 10		
	cctagtccca gggctttgat t	•	. 21
	<210> SEQ ID NO: 11	•	
	<211> LENGTH: 22		
	<212> TYPE: DNA		
	<213> ORGANISM: Artificial Sequence		
	<220> FEATURE:		
	<223> OTHER INFORMATION: Primer		
	<400> SEQUENCE: 11		
	ctgtgctccc actcctgatt tc		22
	<210> SEQ ID NO: 12		
505	<211> LENGTH: 13		

507 <212> TYPE: PRT

<210> 32

<211> 20

<212> DNA

<213> Artificial sequence

needs explanation - see p. 7

<400> 32 gcgcgctagc cacttcggaa

20

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/530,106

DATE: 04/12/2005 TIME: 10:20:32

Input Set : A:\pto.da.txt

on Alpharatron Output Set: N:\CRF4\04122005\J530106.raw

of <220> Feature (NEW RULES):

lence(s) are missing the <220> Feature and associated headings. of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" Unknown". Please explain source of genetic material in <220> to <223> tion (See "Federal Register," 6/01/98, Vol. 63, No. 104,pp.29631-32) c.1.823 of new Rules)

#:32

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/530,106

DATE: 04/12/2005

TIME: 10:20:32

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\04122005\J530106.raw

7 M:270 C: Current Application Number differs, Replaced Current Application No

7 M:271 C: Current Filing Date differs, Replaced Current Filing Date

79 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:32, <213>

ANISM: Artificial sequence

79 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:32, <213>

ANISM: Artificial sequence

79 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:32,Line#:879